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# RAW SEQUENCE LISTING ERROR REPORT

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Application Serial Number: 09   964,05	י נא
Source: OYE	
Date Processed by STIC: 10/08/02	

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

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- 1. EFS-Bio (<a href="http://www.uspto.gov/ebc/efs/downloads/documents.htm">http://www.uspto.gov/ebc/efs/downloads/documents.htm</a>, EFS Submission User Manual ePAVE)
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Revised 01/29/2002



# Does Not Comply Corrected Diskette Needed

OIPE

RAW SEQUENCE LISTING DATE: 10/08/2002 PATENT APPLICATION: US/09/964,059A TIME: 14:15:45

Input Set : N:\EBONY'S\EP.txt

Output Set: N:\CRF4\10082002\I964059A.raw

- 1 <110> APPLICANT: Frudakis, Tony
- 3 <120> TITLE OF INVENTION: Efficient Methods and Apparatus for High-Throughput

## Processing of Gene

- 4 Sequence Data
- 6 <130> FILE REFERENCE: 0201-0001
- 8 <140> CURRENT APPLICATION NUMBER: US 09/964,059A
- 10 <141> CURRENT FILING DATE: 2001-09-26
- 12 <150> PRIOR APPLICATION NUMBER: US 60/274,686
- 13 <151> PRIOR FILING DATE: 2000-03-08
- 15 <160> NUMBER OF SEQ ID NOS: 239

#### **ERRORED SEQUENCES**

- 180 <210> SEQ ID NO: 17 181 <211> LENGTH: 3260
- 182 <212> TYPE: DNA
- 183 <213> ORGANISM: Homo Sapiens
- 185 <400> SEQUENCE: 17 186 agttaatgca cacagtttgg ctagttttgg cttcaaaatt aattaaactg tatcaatgta 187 ttttgaagtg ttaagtcatc tgtatgcttt agctccttct atagatgagg caaatataca 188 aacagattaa actgactttt acagaataat tattetttta cettgtttac atggaaagga 180 189 atcctccatt ttaggatgca cataaaatgc cagcctatgt tgatgacatt gccttaacac 240 190 ttttttttta agtaatttta cagggtagtt aacctgtaaa agaaacagtg gataaacttg 300 191 aaaatgctaa tagcaaaaaa cacttcagcc atggcacata caaccagaag ccaatgatat 360 420 192 ccttcaacta tagaaattag cggtgttttc tgtttattcc tgaagcagga ttccatattc 193 aagccagaaa ttgtcattca acagaaaaaa tcaggtcaaa acaatcaatc acataatgta 480 194 gcaagacaaa agtatgtgct tatgtgaaga aaaacaaaaa caacaaataa ccgaactttt 195 attttcttqa atataatatt qatqqcaaga ttqctaagaq qtcatccctq tatttagttt 600 660 196 agataaaggc ttccagcata gaacactgtt aagaagtaac tgtcaggagc tatgcagaag 197 tgatgagagg caaataatat aaaaactaga aaagcaggtt ttaattttct atagacttta 720 198 ttacacatta ttatgttacg agacaaatgc agataattct taatttatca aatttgtgag 780 199 cttaattaac aaaaatattt gaccctcacc agaaaaacag ataactctaa atctactctg 840 200 aaaatctaat caattqcqaa qtattaccta tttqqaqact atqtattata tcaaaqataa 900 201 agctactatt ctcacagaac atatggggtc attggcagcc aaccaataat gaagtaaata 202 ttctaatatt tgggaaaata ctgagaaaac taataaattg tcctggatat tatttattct 1020 203 tgcctttaca aaagacttac acatccaaat gagattagtt tagaatagag gtttttagtt 1080 204 cagaaaatgt tcaaagtcca atacagtcat ggctaatcag agactagaga acctttataa 1140 205 aggtaagtag gcttgaaaac ccttggaaac tgagcagtct tattttgaac tagcatgttt 1200 206 taatcaaagg tatggaatta atcaaatatc aattaagaat tactggaatg cacactcatg 1260 207 ccaaatgaca actaacatgt tatttcctac tatgatgact ctttgatttg agtcagatgg 1320 208 cataaaaaaa tattgctagc tatacaataa attttactct tctgcttctg ctctctaaag 209 aaaaatctta ttttttcaca taagaagctc atggaatcga atgttaatta aagaaaagat 1440

210 agggtaagta caactggggg aaagacagta cctctaatta cataggaaat ccatgaaaga

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Input Set : N:\EBONY'S\EP.txt

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1560
     211 attaatcatc ataagagaag aatcattttt ccagtagccc cactaccatg aatgatattt
     212 tcatqagcct cqccacctt ctccaatqqa tattqagaac ctatcacagg tttcaaccag
                                                                            1620
     213 ccaatttcca ttccagcttq aaqqqctqct qcatattqct gaaattcctc ctaagaaaaq
                                                                            1680
     214 gaaaaacaaa tttctttttg tagtgaaccg tatgatttaa ttttcagaag cattaaaaac
                                                                            1740
E--> 215 acttcagaat ctaagtgtta taccatgaag agtctcttac aaatgtgtga cttttgtcaa 1700 — 1800
                                                                                    -1860, and so on
E--> 216 cttgtccaga actatagaaa aagtagttat ctacagggta accataaatc ccatctgcct 1760
E--> 217 gagacagtgt tagtgtacaa aatacctgtt gtcctgaaat tattactagt atcacatttc
                                                                            1820
E--> 218 tatctcaaaa ggtatgctta cctggatata aattatactg tcaccctagt tgtccttctg
                                                                            1880
E--> 219 gtgactaatc cttaccaact cccactagtc atataactaa gtttaacatc tattcaaact
E--> 220 ttcagcttgc ctgagtaggc aaactgtacc aatgtttaag ttaccaaaat cagaagtact
E--> 221 tcttttccta ccttggttga ggaaaagaga gtaactccaa ttatactcga ctcctttgcc
                                                                            2060
E--> 222 atggtgtctc gtgggtttat ttcaatagta cctctgctgc caacaaccta acatgaaaaa
                                                                            2120
E--> 223 cagcaattct acagttaaag attactgtaa aatagtgtta aattgtggta aaacattaaa
                                                                            2180
E--> 224 gtggtaaaaa aaaaaaaaag aaaaggaata cttactatca ctcgtcctcc atgtgacaga
                                                                            2240
E--> 225 agactcaagt ctttactaag atttacatta gctaacattt caataattat atcaattcct
                                                                            2300
E--> 226 ttctcaccaa catacttcta tataataaaa gagaaatgta gagtaagata gcaagtgaaa
                                                                            2360
E--> 227 aactgtaaaa tagctactat ctgtacaaga tattatagaa atatgtttca aatgatatat
                                                                            2420
E--> 228 aaatqctaca tctttgagac taataatgca aaattttaaa taatctaatt atataatcac
                                                                            2480
E--> 229 gatgtaattc caaggtacca gccagaacat ctaaactgat aaaaatttgt actaaataca
                                                                            2540
E--> 230 ttgctgtagt gaaataaagt ttgtctggaa ttttcaggtg ctagactcaa cttgagtata
                                                                            2600
E--> 231 aaatacttag ctgaaaattt tctatctgta aaataaactt tcataaagaa acaataaatc
                                                                            2660
E--> 232 aaaagcccca aacccccagg gggctcccat ttttattaat aaacaaaaag caaaagaaga
                                                                            2720
E--> 233 tatcattagc tqttcqqttt tqcatqattt ttqttqtttt agtgcatttg gttttgttct
                                                                            2780
E--> 234 aaatggttta tcatctgttt gatgcactaa ctcttttggg ctcttggatg ttggacgctg
2900
E--> 236 aaatgaatcc ctgtgcccgg gattgcacta ggtaccagga atacaaatac aaacataggg
                                                                            2960
E--> 237 agctcaaaac aaaactagtg agaaagatgg gaaatactac agtcatagct ataaagtaat
                                                                            3020
E--> 238 gggctaagta acacattagc agaaataaat catagaatac agagaaaaaa ggttaaggtt
                                                                            3080
E--> 239 tgattgcctg ccatggtcag ataaagttcc acagagacga tgaactgggc cctcagggat
                                                                            3140
E--> 240 gaataggagt ttcccaagcc aaaagaaagg aaaatgagta aggggaagct agacctgagg
                                                                            3200
E--> 241 ctgagtcagt ctggaccaaa gaaacagaaa agcaaagatg gaggggactg agaacacaag
                                                                            3260
                                             Ms in sequence
P. 6 for exp bonetion
     340 <210> SEQ ID NO: 20
     341 <211> LENGTH: 384
                                                                        The type of errors shown exist throughout
     342 <212> TYPE: DNA
                                                                       A the Sequence Listing. Please check subseque
     343 <213> ORGANISM: Homo Sapiens
                                                                        sequences for similar errors.
     345 <400> SEQUENCE: 20
     346 acaqacaqaq ataqacaqaq qcaqaqtcaq qqaqaqqcaq agaaaqaaag agaacaaqaa
                                                                            60
     347 agettaaaga tagtecaaac geaaagetgt etttaaaaaa tgeataetet attaetggea
     348 acaaagtttt ataatctata cattttatga accactaatc cttaatttat tcaagatcac
     349 aacaggggac tcatattata gagtcaagta aatatcatta ccaacatttt atttaacagt
                                                                            240
     350 ttgtcctcct taattacatg gagaatgata tagtgactcc ttcatgcctt tttttctcct
                                                                            300
     351 taacaagcca tatgcaggaa agtttccatg ctgcgcaaac ataaaagaaa gttatatttc
E--> 352 attectains gaaaactgaa aage
     815 <210> SEQ ID NO: 63
     816 <211> LENGTH: 1300
                                            Some ever
     817 <212> TYPE: DNA
     818 <213> ORGANISM: Homo Sapiens
     820 <400> SEQUENCE: 63
E--> 821 mnnattcagt accaaattta ctgtggaaac agccagtaga gaatacaaga aaatgttcaa 60
```

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```
822 acaggcaagt aaataagtgt cttgtacctt aatgataaat ggtagtagta tagccattta
                                                                             120
     823 taatqqcatt aatqattqqt ttaatttaac ataatttata agctattqaa qtatqqaaaa
                                                                             180
     824 ttataagcat atatattagg ttattaggac tcataaattt atgttattta cttccagttt
                                                                             240
     825 gtgagatgac ttgaattttt catgtttcct attctttact tccatagaca tggatgqata
                                                                             300
     826 atatgggaag agctggtgag atggaactca agcccttcaa tggagaagat tatacatgta
                                                                             360
     827 tcacctttca gcctgatttg tctaagttta aaatgcaaag cctggacaaa gatattgttg
                                                                             420
     828 cactaatggt cagaagagca tatgatattg ctggatccac caaagatgtc aaagtctttc
                                                                             480
     829 ttaatggaaa taaactgcca tgagtatttt cctggatgtt aaggataata agggattttg
                                                                             540
     830 taatcattgt caagtgcaaa attgaatttt ttcccctccc atatgttttt gtttgtttgt
     831 ttgtttgttt gtttgagaca gagtctcaca ctgttgcccg ggctggagtg cagtggcacg
                                                                             660
     832 attecegete accgeaacet ceacetecea ggtteaegea atteteetge eteageetee
                                                                             720
     833 caagtagetg ggattacagg tgcctgccac cacacctggc taattttttg tatttttagt
                                                                             780
     834 agagacaggt ttcactatgt tggccaggct ggtctcgaac accagacctc atgatccacc
                                                                             840
     835 cgtcttggcc tcccaaagtg ctgggattac aggcatgagc cactgcacct ggcccaacca
                                                                             900
     836 tatgtatttt cttaccactt ctcacatatg ttcttgaaaa gagaatggta tgccacattt
                                                                             960
     837 tttaatcagc tcattttaaa cttaccgaag gaatttcttt ctcaaagaaa cacctaaaat
     838 aaatatttca tgtccttttt ttattttcct ttttctttct tttcttgata acctcgctgt
                                                                             1080
     839 gtcacccagg ctggagtaca gtgatgcaat cacggctcac tacagcctgg acctcccagg
                                                                             1140
     840 ctcaagcgat catcccacct cagcttctqq agtaqctgga aatqcagqca qcaccaccat
                                                                             1200
     841 gcccagctaa ttttttttt tctttttaat agaggtgggg atctcactat gttgcccagg
     842 ctggtcttga actcctgggc tcaagtgatc cacccacctc
     1579 <210> SEQ ID NO: 146
     1580 <211> LENGTH: 41
                                                                       Some errors as Seq., 20
     1581 <212> TYPE: DNA
     1582 <213> ORGANISM: Homo Sapiens
     1584 <400> SEQUENCE: 146
E--> 1585 agggggtaga ttttaaaaat ncatgttaat gttatttact n
     1587 <210> SEQ ID NO: 147
     1588 <211> LENGTH: 41
     1589 <212> TYPE: DNA
     1590 <213> ORGANISM: Homo Sapiens
     1592 <400> SEQUENCE: 147
E--> 1593 agggggtaga ttttaaaaat ncatgttaat gttatttact n
                                                            41
     1595 <210> SEQ ID NO: 148
     1596 <211> LENGTH: 41
     1597 <212> TYPE: DNA
     1598 <213> ORGANISM: Homo Sapiens
     1600 <400> SEQUENCE: 148
E--> 1601 aggtgtaaga ttttaaaaat acatgttaat gttatttact n
                                                            41
     1603 <210> SEQ ID NO: 149
     1604 <211> LENGTH: 41
     1605 <212> TYPE: DNA
     1606 <213> ORGANISM: Homo Sapiens
     1608 <400> SEQUENCE: 149
E--> 1609 aggggtanga tttcaaaaat acatgttaat gttatttact n
     1611 <210> SEQ ID NO: 150
     1612 <211> LENGTH: 41
     1613 <212> TYPE: DNA
```

1614 <213> ORGANISM: Homo Sapiens

RAW SEQUENCE LISTING

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```
E--> 1617 aggggtanga ttttaaaaat acatgttaat gttatttact n
                                                             41
     1619 <210> SEQ ID NO: 151
     1620 <211> LENGTH: 41
     1621 <212> TYPE: DNA
     1622 <213> ORGANISM: Homo Sapiens
     1624 <400> SEQUENCE: 151
E--> 1625 aggggtaaga ttttaaaaat acatgttaat gttatttact n
                                                             41
     1627 <210> SEQ ID NO: 152
     1628 <211> LENGTH: 41
     1629 <212> TYPE: DNA
     1630 <213> ORGANISM: Homo Sapiens
     1632 <400> SEQUENCE: 152
E--> 1633 aggggtaaga ttttaaaaaat acatgttaat gttatttact n
                                                             41
     1687 <210> SEQ ID NO: 158
     1688 <211> LENGTH: 11
     1689 <212> TYPE: DNA
     1690 <213> ORGANISM: Homo Sapiens
     1692 <400> SEQUENCE: 158
E--> 1693 cntctgagtt a
                                 11
     1695 <210> SEQ ID NO: 159
     1696 <211> LENGTH: 11
     1697 <212> TYPE: DNA
     1698 <213> ORGANISM: Homo Sapiens
     1700 <400> SEQUENCE: 159
E--> 1701 cntctgagtt a
                                 11
     1703 <210> SEQ ID NO: 160
     1704 <211> LENGTH: 11
     1705 <212> TYPE: DNA
     1706 <213> ORGANISM: Homo Sapiens
     1708 <400> SEQUENCE: 160
E--> 1709 cntttgagtt a
                                 11
     1711 <210> SEQ ID NO: 161
     1712 <211> LENGTH: 11
     1713 <212> TYPE: DNA
     1714 <213> ORGANISM: Homo Sapiens
     1716 <400> SEQUENCE: 161
E--> 1717 cntctgagtt a
                                 11
     1719 <210> SEQ ID NO: 162
     1720 <211> LENGTH: 11
     1721 <212> TYPE: DNA
     1722 <213> ORGANISM: Homo Sapiens
     1724 <400> SEQUENCE: 162
E--> 1725 cntttgagtt a
                                 11
```

1727 <210> SEQ ID NO: 163 1728 <211> LENGTH: 11 1729 <212> TYPE: DNA

1732 <400> SEQUENCE: 163

1730 <213> ORGANISM: Homo Sapiens

RAW SEQUENCE LISTING DATE: 10/08/2002 PATENT APPLICATION: US/09/964,059A TIME: 14:15:45

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```
Some errors as M Seq, 20
E--> 1733 cntctgagtt a
                                 11
     1735 <210> SEQ ID NO: 164
     1736 <211> LENGTH: 11
     1737 <212> TYPE: DNA
     1738 <213> ORGANISM: Homo Sapiens
     1740 <400> SEQUENCE: 164
E--> 1741 cntctgagtt a
                                 11
     1743 <210> SEQ ID NO: 165
     1744 <211> LENGTH: 11
     1745 <212> TYPE: DNA
     1746 <213> ORGANISM: Homo Sapiens
     1748 <400> SEQUENCE: 165
E--> 1749 cntctgagtt a
                                 11
     1751 <210> SEQ ID NO: 166
     1752 <211> LENGTH: 11
     1753 <212> TYPE: DNA
     1754 <213> ORGANISM: Homo Sapiens
     1756 <400> SEQUENCE: 166
E--> 1757 cntctgagtt a
                                 11
     1759 <210> SEQ ID NO: 167
     1760 <211> LENGTH: 11
     1761 <212> TYPE: DNA
     1762 <213> ORGANISM: Homo Sapiens
     1764 <400> SEQUENCE: 167
E--> 1765 cntctgagtt a
     1767 <210> SEQ ID NO: 168
     1768 <211> LENGTH: 11
     1769 <212> TYPE: DNA
     1770 <213> ORGANISM: Homo Sapiens
     1772 <400> SEQUENCE: 168
E--> 1773 cntttgagtt a
     2330 <210> SEQ ID NO: 238
     2331 <211> LENGTH: (260) - Found 200
     2332 <212> TYPE: DNA
     2333 <213> ORGANISM: Homo Sapiens
     2335 <400> SEQUENCE: 238
     2336 etcaggtece acagcaacaa tateatteaa actgeaatta aaacatacae acataatata 60
E--> 2337 taaggtgaag gtattgaaca ttacaggatt attaactggc attcctcact gtctattcct (180) \rightarrow 120
E--> 2338 aaaatcaaga tgtgggatgg agccttcgtg ctagctataa tggaacacaa ttaatatgaa (240) \rightarrow 180
                                         (260) - 200
E--> 2339 attagtcctg ccgatacaat
```

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### Use of n's or Xaa's (NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing. Use of  $\langle 220 \rangle$  to  $\langle 223 \rangle$  is MANDATORY if n's or Xaa's are present. in  $\langle 220 \rangle$  to  $\langle 223 \rangle$  section, please explain location of n or Xaa, and which residue n or Xaa represents.

```
Seq#:20; N Pos. 369
Seq#:63; N Pos. 1,2,3
Seq#:146; N Pos. 21,41
Seq#:147; N Pos. 21,41
Seq#:148; N Pos. 41
Seq#:149; N Pos. 8,41
Seq#:150; N Pos. 8,41
Seq#:151; N Pos. 41
Seq#:152; N Pos. 41
Seq#:158; N Pos. 2
Seq#:159; N Pos. 2
Seq#:160; N Pos. 2
Seq#:161; N Pos. 2
Seq#:162; N Pos. 2
Seq#:163; N Pos. 2
Seq#:164; N Pos. 2
Seq#:165; N Pos. 2
Seq#:166; N Pos. 2
Seq#:167; N Pos. 2
Seq#:168; N Pos. 2
Seq#:169; N Pos. 2
Seq#:170; N Pos. 2
Seq#:171; N Pos. 2
Seq#:172; N Pos. 2
Seq#:173; N Pos. 2
Seq#:174; N Pos. 2
Seq#:175; N Pos. 2
Seq#:177; N Pos. 4
Seq#:178; N Pos. 4
Seq#:179; N Pos. 4
Seq#:180; N Pos. 4
Seq#:181; N Pos. 4
Seq#:182; N Pos. 1,2
Seq#:183; N Pos. 1,2,3,4,5,6
Seq#:184; N Pos. 2
Seq#:185; N Pos. 2
Seq#:186; N Pos. 2
Seq#:187; N Pos. 2
Seq#:188; N Pos. 2
Seq#:189; N Pos. 4
Seq#:190; N Pos. 2
Seq#:191; N Pos. 2
Seq#:192; N Pos. 2
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VARIABLE LOCATION SUMMARY

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Input Set : N:\EBONY'S\EP.txt

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```
Seq#:193; N Pos. 2
Seq#:194; N Pos. 4
Seq#:195; N Pos. 4
Seq#:199; N Pos. 2
Seq#:200; N Pos. 4
Seq#:204; N Pos. 4
Seq#:206; N Pos. 5
Seq#:208; N Pos. 4
Seq#:209; N Pos. 2
Seq#:210; N Pos. 2
Seq#:212; N Pos. 2
Seq#:213; N Pos. 2
Seq#:214; N Pos. 1,2,10
Seq#:218; N Pos. 2
Seq#:219; N Pos. 2
Seq#:220; N Pos. 96,120,148,149,150,175
Seq#:235; N Pos. 9,10,12
Seq#:236; N Pos. 9,10,11,12,13
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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/964,059A

TIME: 14:15:46

Input Set : N:\EBONY'S\EP.txt

Output Set: N:\CRF4\10082002\I964059A.raw

L:215 M:254 E: No. of Bases conflict, LENGTH:Input:1700 Counted:1800 SEQ:17 M:254 Repeated in SeqNo=17 L:241 M:252 E: No. of Seq. differs, <211> LENGTH:Input:3260 Found:3360 SEQ:17 L:352 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:20 L:821 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:63 L:1585 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:146 L:1593 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:147 L:1601 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:148 L:1609 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:149 L:1617 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:150 L:1625 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:151 L:1633 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:152 L:1693 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:158 L:1701 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:159 L:1709 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:160 L:1717 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:161 L:1725 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:162 L:1733 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:163 L:1741 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:164 L:1749 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:165 L:1757 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:166 L:1765 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:167 L:1773 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:168 L:1781 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:169 L:1789 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:170 L:1797 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:171 L:1805 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:172 L:1813 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:173 L:1821 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:174 L:1829 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:175 L:1845 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:177 L:1853 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:178 L:1861 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:179 L:1869 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:180 L:1877 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:181 L:1885 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:182 L:1893 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:183 L:1901 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:184 L:1909 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:185 L:1917 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:186 L:1925 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:187 L:1933 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:188 L:1941 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:189 L:1949 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:190 L:1957 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:191 L:1965 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:192 L:1973 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:193 L:1981 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:194

VERIFICATION SUMMARY

DATE: 10/08/2002 TIME: 14:15:46

PATENT APPLICATION: US/09/964,059A

Input Set : N:\EBONY'S\EP.txt

Output Set: N:\CRF4\10082002\I964059A.raw

L:1989 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:195 L:2021 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:199 L:2029 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:200 L:2061 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:204 L:2077 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:204 L:2337 M:254 E: No. of Bases conflict, LENGTH:Input:180 Counted:120 SEQ:238 M:254 Repeated in SeqNo=238

L:2339 M:252 E: No. of Seq. differs, <211> LENGTH:Input:260 Found:200 SEQ:238